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SEQUENCE LISTING

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<110> ANDREEV, et al.

<120> METHODS FOR IDENTIFYING MODULATORS OF ACTIVE KIT TYROSINE KINASE RECEPTOR

<130> 30694/39618A

<150> US 60/526,930

<151> 2003-12-04

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<170> PatentIn version 3.2

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Leu Gly Ser Ser Pro Tyr Pro Gly Met Pro Val Asp Ser Lys Phe Tyr
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Val Val Glu Lys Gly Phe Ile Asn Ile Phe Pro Val Lys Asn Thr Thr	
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Val Phe Val Thr Asp Gly Glu Asn Val Asp Leu Val Val Glu Phe Glu	
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gcc tac cct aaa cct gaa cac cag cag tgg atc tac atg aac agg acg	1112
Ala Tyr Pro Lys Pro Glu His Gln Gln Trp Ile Tyr Met Asn Arg Thr	
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Pro Thr Asn Arg Gly Glu Asp Tyr Val Lys Ser Asp Asn Gln Ser Asn	
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Ile Arg Tyr Val Asn Glu Leu Arg Leu Thr Arg Leu Lys Gly Thr Glu	
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Gly Gly Thr Tyr Thr Phe Leu Val Ser Asn Ser Asp Val Ser Ala Ser	
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Val Thr Phe Asp Val Tyr Val Asn Thr Lys Pro Glu Ile Leu Thr Tyr	
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Asp Arg Leu Met Asn Gly Arg Leu Gln Cys Val Ala Ala Gly Phe Pro	
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Glu Pro Thr Ile Asp Trp Tyr Phe Cys Thr Gly Ala Glu Gln Arg Cys	
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Thr Val Pro Val Pro Val Asp Val Gln Ile Gln Asn Ala Ser Val	
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Ser Pro Phe Gly Lys Leu Val Val Gln Ser Ser Ile Asp Ser Ser Val	
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Phe Arg His Asn Gly Thr Val Glu Cys Lys Ala Ser Asn Ala Val Gly	
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Lys Ser Ser Ala Phe Phe Asn Phe Ala Phe Lys Gly Asn Ser Lys Glu	
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Gln Ile Gln Pro His Thr Leu Phe Thr Pro Leu Leu Ile Gly Phe Val	
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Val Thr Ala Gly Leu Met Gly Ile Ile Val Met Val Leu Ala Tyr Lys	
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gcg	tgt	acc	gtg	gga	ggg	ccc	acc	ctg	gtc	att	aca	gaa	tac	tgt	tgc	2072
Ala	Cys	Thr	Val	Gly	Gly	Pro	Thr	Leu	Val	Ile	Thr	Glu	Tyr	Cys	Cys	
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Phe	Ser	Lys	Gln	Glu	Glu	Gln	Ala	Asp	Ala	Ala	Leu	Tyr	Lys	Asn	Leu	
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Ile	Met	Glu	Asp	Asp	Glu	Leu	Ala	Leu	Asp	Leu	Glu	Asp	Leu	Leu	Ser	
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Phe	Ser	Tyr	Gln	Val	Ala	Lys	Gly	Met	Ala	Phe	Leu	Ala	Ser	Lys	Asn	
		775					780					785				
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Cys	Ile	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Ile	Leu	Leu	Thr	His	Gly	
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Val Trp Ser Tyr Gly Ile Phe Leu Trp Glu Leu Phe Ser Leu Gly Ser	
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Ser Lys His Ile Tyr Ser Asn Leu Ala Asn Cys Asn Pro Asn Pro Glu	
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Asn Pro Val Val Val Asp His Ser Val Arg Val Asn Ser Val Gly Ser	
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agc acc tct tcc aca cag cct ctc ctc gtg cat gag gac gcc tga	2981
Ser Thr Ser Ser Thr Gln Pro Leu Leu Val His Glu Asp Ala	
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35 40 45

Glu Ala Gly Asp Thr Ile Arg Leu Thr Cys Thr Asp Pro Ala Phe Val
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Lys Trp Thr Phe Glu Ile Leu Asp Val Arg Ile Glu Asn Lys Gln Ser
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Glu Trp Ile Arg Glu Lys Ala Glu Ala Thr His Thr Gly Lys Tyr Thr
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Cys Val Ser Gly Ser Gly Leu Arg Ser Ser Ile Tyr Val Phe Val Arg
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Asp Pro Ala Val Leu Phe Leu Val Gly Leu Pro Leu Phe Gly Lys Glu
115 120 125

Asp Asn Asp Ala Leu Val Arg Cys Pro Leu Thr Asp Pro Gln Val Ser
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Asn Tyr Ser Leu Ile Glu Cys Asp Gly Lys Ser Leu Pro Thr Asp Leu
145 150 155 160

Lys Phe Val Pro Asn Pro Lys Ala Gly Ile Thr Ile Lys Asn Val Lys
165 170 175

Arg Ala Tyr His Arg Leu Cys Ile Arg Cys Ala Ala Gln Arg Glu Gly
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Lys Trp Met Arg Ser Asp Lys Phe Thr Leu Lys Val Arg Ala Ala Ile
 195 200 205
 Lys Ala Ile Pro Val Val Ser Val Pro Glu Thr Ser His Leu Leu Lys
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 Glu Gly Asp Thr Phe Thr Val Ile Cys Thr Ile Lys Asp Val Ser Thr
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 Lys Asn Thr Thr Val Phe Val Thr Asp Gly Glu Asn Val Asp Leu Val
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 Met Asn Arg Thr Pro Thr Asn Arg Gly Glu Asp Tyr Val Lys Ser Asp
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 Ala Gly Phe Pro Glu Pro Thr Ile Asp Trp Tyr Phe Cys Thr Gly Ala
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Glu Gln Arg Cys Thr Val Pro Val Pro Pro Val Asp Val Gln Ile Gln
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 Ile Gly Phe Val Val Thr Ala Gly Leu Met Gly Ile Ile Val Met Val
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 Ser Glu Leu Lys Val Leu Ser Tyr Leu Gly Asn His Met Asn Ile Val
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 Asp Ser Phe Ile Phe Ser Lys Gln Glu Glu Gln Ala Asp Ala Ala Leu
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Glu	Tyr	Met	Asp	Met	Lys	Pro	Gly	Val	Ser	Tyr	Val	Val	Pro	Thr	Lys	
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Ser Val Gly Ser Ser Thr Ser Ser Thr Gln Pro Leu Leu Val His Glu
965 970 975

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